

OIPE

RAW SEQUENCE LISTING DATE: 09/18/2001 PATENT APPLICATION: US/09/942,407 TIME: 16:20:49

Input Set : $D:\30062-20049.txt$

Output Set: N:\CRF3\09182001\I942407.raw



			out	Puc	Dec.	11.	(0111 0	. (03 -		- \						r¢ RED
4	<110>	APPLI	CANT:	Kos	an B	iosc	ienc	es,	Inc.						7	ı
5		Santi														
6		Peck,									•		اں دے	Ail II		
7		Dayem		_								į				
8		Kealey														
	<120>	TITLE	OF I	NVEN	TION	ı: IS	OLAT	ED G	ENE	FOR	METH	IYLMA	LONY	L CC	A EPI	MERASE AND USES
THEREOF	1221															
	<130>	FILE P	REFER	RENCE	: 30	062-	2004	9.10)							•
C> 15										942,	407					
C> 16										-						
		PRIOR								9,13	16					
		PRIOR														
		PRIOR							0/16	1,41	. 4					
		PRIOR														
		PRIOR							0/16	1,70	3					
		PRIOR														
		PRIOR							0/20	6,08	32					
		PRIOR							·							
		NUMBE														
		SOFTW						awof	Vers	ion	4.0					
		SEQ I														•
35	<211>	LENGT	H: 44	17												
		TYPE:														•
37	<213>	ORGAN	ISM:	Arti	ifici	ial S	Seque	ence								
39	<220>	FEATU	RE:								1					
40	<223>	OTHER	INFO	RMAT	ON:	: Isc	olate	ed ar	nd re	comb	oinar	nt fo	orm o	of th	ne ful	س 11
41		epime:	rase	gene	e sec	quenc	ce									
43	<221>	NAME/	KEY:	CDS												
44	<222>	LOCAT	: NOI	(1)	(4	144)										
		SEQUE:														
		gt aat														48
48	Met S	Ser Asn	Glu	Asp	Leu	Phe	Ile	Cys		Asp	His	Val	Ala		Ala	
	1			5					10					15		
51	tgc c	ccc gac	gcc	gac	gag	gct	tcc	aag	tac	tac	cag	gag	acc	ttc	ggc	96
52	Cys F	ro Asp		Asp	Glu	Ala	Ser		Tyr	Tyr	Gln	Glu		Phe	Gly	
53			20					25					30			
55	tgg c	cat gag	ctc	cac	cgc	gag	gag	aac	ccg	gag	cag	gga	gtc	gtc	gag	144
56	Trp F	is Glu	Leu	His	Arg	Glu	Glu	Asn	Pro	Glu	Gln	Gly	Val	Val	Glu	
57		35					40					45				
59	atc a	atg atg	gcc	ccg	gct	gcg	aag	ctg	acc	gag	cac	atg	acc	cag	gtt	192
60	Ile M	Met Met	Ala	Pro	Ala		Lys	Leu	Thr	Glu		Met	Thr	Gln	Val	
61		50				55					60					2.4.2
		gtc atg														240
		/al Met	Ala	Pro		Asn	Asp	Glu	Ser		Val	Ala	Lys	Trp		
	65				70					75			4		80	200
. 67	gcc a	aag cac	aat	ggt	cgc	gcc	gga	ctg	cac	cac	atg	gca	rgg	cgt	gtc	288
68	7 1 = 1	Lys His	λcn	C 1 37	7 200	Δla	C137	1.011	Hi C	U 1 C	MOT	$\Delta I \Delta$	מאיוי	Ara	vai	
69		ays mis	HSII	85	AIG	пια	GLY	пси	90	птъ	Mec	AIG	111	95	, u _	



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71 gat gac atc gac gcc gtc agc gcc acc ctg cgc gag cgc ggc gtg cag 72 Asp Asp Ile Asp Ala Val Ser Ala Thr Leu Arg Glu Arg Gly Val Gln 73 100 105 110	336
75 ctg ctg tat gac gag ccc aag ctc ggc acc ggc ggc aac cgc atc aac	384
76 Leu Leu Tyr Asp Glu Pro Lys Leu Gly Thr Gly Gly Asn Arg Ile Asn	
77 115 120 125	
79 ttc atg cat ccc aag tcg ggc aag ggc gtg ctc atc gag ctc acc cag	432
80 Phe Met His Pro Lys Ser Gly Lys Gly Val Leu Ile Glu Leu Thr Gln	
81 130 135 140	4 4 7
83 tac ccg aag aac tga	447
84 Tyr Pro Lys Asn	
85 145	
88 <210> SEQ ID NO: 2 89 <211> LENGTH: 148	
90 <212> TYPE: PRT	
91 <213> ORGANISM: Artificial Sequence	
93 <220> FEATURE:	
94 <223> OTHER INFORMATION: Deduced amino acid sequence of the epimeras	e gene
95 sequence	
97 <400> SEQUENCE: 2	
on wet dem has die hes tou Dhe the due the hes Wel his wor his	
98 Met Ser Asn Glu Asp Leu Phe Ile Cys Ile Asp His Val Ala Tyr Ala	
99 1 5 10 15	
99 1 5 10 15 100 Cys Pro Asp Ala Asp Glu Ala Ser Lys Tyr Tyr Gln Glu Thr Phe Gly 101 20 25 30	
99 1 5 10 15 100 Cys Pro Asp Ala Asp Glu Ala Ser Lys Tyr Tyr Gln Glu Thr Phe Gly	
99 1 5 10 15 100 Cys Pro Asp Ala Asp Glu Ala Ser Lys Tyr Tyr Gln Glu Thr Phe Gly 101 20 25 30 102 Trp His Glu Leu His Arg Glu Glu Asn Pro Glu Gln Gly Val Val Glu 103 35 40 45	
99 1 5 10 15 100 Cys Pro Asp Ala Asp Glu Ala Ser Lys Tyr Tyr Gln Glu Thr Phe Gly 101 20 25 30 102 Trp His Glu Leu His Arg Glu Glu Asn Pro Glu Gln Gly Val Val Glu 103 35 40 45 104 Ile Met Met Ala Pro Ala Ala Lys Leu Thr Glu His Met Thr Gln Val 105 50 55 60	
99 1 5 10 15 100 Cys Pro Asp Ala Asp Glu Ala Ser Lys Tyr Tyr Gln Glu Thr Phe Gly 101 20 25 30 102 Trp His Glu Leu His Arg Glu Glu Asn Pro Glu Gln Gly Val Val Glu 103 35 40 45 104 Ile Met Met Ala Pro Ala Ala Lys Leu Thr Glu His Met Thr Gln Val	
99 1 5 10 15 15 10 15 15 10 15 10 Cys Pro Asp Ala Asp Glu Ala Ser Lys Tyr Tyr Gln Glu Thr Phe Gly 101 20 25 30 30 102 Trp His Glu Leu His Arg Glu Glu Asn Pro Glu Gln Gly Val Val Glu 103 35 40 45 104 Ile Met Met Ala Pro Ala Ala Lys Leu Thr Glu His Met Thr Gln Val 105 50 50 55 60 106 Gln Val Met Ala Pro Leu Asn Asp Glu Ser Thr Val Ala Lys Trp Leu 107 65 70 70 75 80	
99 1 5 10 15 100 Cys Pro Asp Ala Asp Glu Ala Ser Lys Tyr Tyr Gln Glu Thr Phe Gly 101 20 20 25 30 102 Trp His Glu Leu His Arg Glu Glu Asn Pro Glu Gln Gly Val Val Glu 103 35 40 45 104 Ile Met Met Ala Pro Ala Ala Lys Leu Thr Glu His Met Thr Gln Val 105 50 55 60 106 Gln Val Met Ala Pro Leu Asn Asp Glu Ser Thr Val Ala Lys Trp Leu 107 65 70 75 80 108 Ala Lys His Asn Gly Arg Ala Gly Leu His His Met Ala Trp Arg Val	
99 1 5 10	
99 1 5 5 10 15 15 15 10 15 15 100 Cys Pro Asp Ala Asp Glu Ala Ser Lys Tyr Tyr Gln Glu Thr Phe Gly 101 20 25 30 30 102 Trp His Glu Leu His Arg Glu Glu Asn Pro Glu Gln Gly Val Val Glu 103 35 40 45 45 104 Ile Met Met Ala Pro Ala Ala Lys Leu Thr Glu His Met Thr Gln Val 105 50 50 55 60 106 Gln Val Met Ala Pro Leu Asn Asp Glu Ser Thr Val Ala Lys Trp Leu 107 65 70 70 75 80 80 108 Ala Lys His Asn Gly Arg Ala Gly Leu His His Met Ala Trp Arg Val 109 85 90 95 110 Asp Asp Ile Asp Ala Val Ser Ala Thr Leu Arg Glu Arg Gly Val Gln	
99 1 5 5 10 15 15 15 10 15 15 100 Cys Pro Asp Ala Asp Glu Ala Ser Lys Tyr Tyr Gln Glu Thr Phe Gly 101 20 25 30 30 102 Trp His Glu Leu His Arg Glu Glu Asn Pro Glu Gln Gly Val Val Glu 103 35 40 45 45 104 Ile Met Met Ala Pro Ala Ala Lys Leu Thr Glu His Met Thr Gln Val 105 50 50 55 60 106 Gln Val Met Ala Pro Leu Asn Asp Glu Ser Thr Val Ala Lys Trp Leu 107 65 70 70 75 80 80 108 Ala Lys His Asn Gly Arg Ala Gly Leu His His Met Ala Trp Arg Val 109 85 90 95 110 Asp Asp Ile Asp Ala Val Ser Ala Thr Leu Arg Glu Arg Gly Val Gln 111 100 105 105	
99 1 5 5 6 10 Asp Pro Asp Ala Asp Glu Ala Ser Lys Tyr Tyr Gln Glu Thr Phe Gly 101 20 20 25 30 102 Trp His Glu Leu His Arg Glu Glu Asn Pro Glu Gln Gly Val Val Glu 103 35 40 45 104 Ile Met Met Ala Pro Ala Ala Lys Leu Thr Glu His Met Thr Gln Val 105 50 50 55 60 106 Gln Val Met Ala Pro Leu Asn Asp Glu Ser Thr Val Ala Lys Trp Leu 107 65 70 70 75 80 108 Ala Lys His Asn Gly Arg Ala Gly Leu His His Met Ala Trp Arg Val 109 85 90 95 110 Asp Asp Ile Asp Ala Val Ser Ala Thr Leu Arg Glu Arg Gly Val Gln 111 100 105 Leu Gly Thr Gly Gly Asn Arg Ile Asn	
99 1 5 5 10 15 15 10 15 15 10 Cys Pro Asp Ala Asp Glu Ala Ser Lys Tyr Tyr Gln Glu Thr Phe Gly 101 20 25 30 30 102 Trp His Glu Leu His Arg Glu Glu Asn Pro Glu Gln Gly Val Val Glu 103 35 40 45 104 Ile Met Met Ala Pro Ala Ala Lys Leu Thr Glu His Met Thr Gln Val 105 50 50 55 60 106 Gln Val Met Ala Pro Leu Asn Asp Glu Ser Thr Val Ala Lys Trp Leu 107 65 70 70 75 80 80 108 Ala Lys His Asn Gly Arg Ala Gly Leu His His Met Ala Trp Arg Val 109 85 90 95 110 Asp Asp Ile Asp Ala Val Ser Ala Thr Leu Arg Glu Arg Gly Val Gln 111 100 105 110 112 Leu Leu Tyr Asp Glu Pro Lys Leu Gly Thr Gly Gly Asn Arg Ile Asn 113 115 120 125	
99 1 5 5 10 10 15 15 10 15 10 15 10 10 15 10 10 Cys Pro Asp Ala Asp Glu Ala Ser Lys Tyr Tyr Gln Glu Thr Phe Gly 101 20 20 25 30 30 102 Trp His Glu Leu His Arg Glu Glu Asn Pro Glu Gln Gly Val Val Glu 103 35 40 45 104 116 Met Met Ala Pro Ala Ala Lys Leu Thr Glu His Met Thr Gln Val 50 50 55 60 106 Gln Val Met Ala Pro Leu Asn Asp Glu Ser Thr Val Ala Lys Trp Leu 107 65 70 70 75 80 80 108 Ala Lys His Asn Gly Arg Ala Gly Leu His His Met Ala Trp Arg Val 109 85 90 95 110 Asp Asp Ile Asp Ala Val Ser Ala Thr Leu Arg Glu Arg Gly Val Gln 111 100 100 105 110 110 111 115 115 116 Asp Ala Val Pro Lys Leu Gly Thr Gly Gly Asn Arg Ile Asn 113 115 120 120 125 114 Phe Met His Pro Lys Ser Gly Lys Gly Val Leu Ile Glu Leu Thr Gln	
99 1	
99 1 5 5 10 10 15 15 10 15 10 15 10 10 15 10 10 Cys Pro Asp Ala Asp Glu Ala Ser Lys Tyr Tyr Gln Glu Thr Phe Gly 101 20 20 25 30 30 102 Trp His Glu Leu His Arg Glu Glu Asn Pro Glu Gln Gly Val Val Glu 103 35 40 45 104 116 Met Met Ala Pro Ala Ala Lys Leu Thr Glu His Met Thr Gln Val 50 50 55 60 106 Gln Val Met Ala Pro Leu Asn Asp Glu Ser Thr Val Ala Lys Trp Leu 107 65 70 70 75 80 80 108 Ala Lys His Asn Gly Arg Ala Gly Leu His His Met Ala Trp Arg Val 109 85 90 95 110 Asp Asp Ile Asp Ala Val Ser Ala Thr Leu Arg Glu Arg Gly Val Gln 111 100 100 105 110 110 111 115 115 116 Asp Ala Val Pro Lys Leu Gly Thr Gly Gly Asn Arg Ile Asn 113 115 120 120 125 114 Phe Met His Pro Lys Ser Gly Lys Gly Val Leu Ile Glu Leu Thr Gln	





VERIFICATION SUMMARY
PATENT APPLICATION: US/09/942,407

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L:15 M:270 C: Current Application Number differs, Replaced Current Application Number

L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date